



SEQUENCE LISTING

<110> Vert, Patrick J.
Brewer, George J.
Vilma, Yuzbasiyan-Gurkan
Schall, William D.

<120> DNA ENCODING CANINE VON WILLEBRAND FACTOR AND METHODS
OF USE

<130> UMW-1326CPPCS

<150> 09/132,652

<151> 1998-08-11

<150> PCT/US99/18153

<151> 1999-08-10

<160> 29

<170> PatentIn Ver. 2.0

<210> 1

<211> 8402

<212> DNA

<213> Canis familiaris

<400> 1

```
cattaaaaagg tectgggttg gagttttttt ttgggaccag cactccatgt tcaagggpaa 60
acagggggaa attaggatca atcttttttt ttcttttttt taaaaaaaat aattcttccc 120
actttgaaa cggacagtag tacataccag tagctctctg cgaggacggt gatcactaat 180
cattttctt gcttctgggc agatgagtc taccagaatt gtgagggtgt tpttggctct 240
ggccctcctc ttgcccaggga aactttgtac aaaagggact gttggaaggt catgagtggc 300
cggatgacg ctctctggag gtaacttat caacaccttt gatgagatc ttttctctt 360
tgggggagat tggagtlacc tcttgggttg ggaactccag gaacactcga tttctttat 420
ggggggatc caaatgaca aaagagtgag cttctctgtg tatctgggag aatttttga 480
tattttttt ttttttctg gtacatctt ggaatttacc caaagtatc ttttttcta 540
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 600
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 660
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 720
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 780
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 840
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 900
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 960
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 1020
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 1080
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 1140
ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt ttttttttt 1200
```

ataaggggag	gaactgcaag	tggattcggg	gttccggggg	aggtaactgg	tgaagctgta	1740
cccccgcctar	gaggggaaag	cgtgggggag	tgggggggaa	tacaaaggga	acggggggga	1800
ggatttctta	aggggagag	gactggggga	gactctggg	gacttattcg	ggaaagcctg	1860
gaagctggtt	aggggctggg	agaaactgca	gagggagga	gaggtatcct	ggagcctcga	1920
cccgaggacag	gaaaggtttg	eggaggaggc	gtgggggctg	ctgaggtcct	cgaagttcga	1980
gccttgaaac	gaggggtggg	gtcctcagcc	ctaatgtgag	aatgactctc	acgaactctg	2040
ctactgctcc	gaaggcagag	actgtctctg	cagggccttg	gcaatctacg	cggcaggcctg	2100
ggcccgaggg	ggagtgcaca	tccggtgggg	ggagccgggg	ttctgtggcg	tgaagtgccc	2160
ccaggggccag	gtgtacctgc	agtgtgggac	cccttgcaac	atgaactgtc	tctccctctc	2220
ttaccgggag	gaggactgca	atgaggctctg	cttggaaaag	tgtttctccc	cccccagggt	2280
gtacctggat	gagaggggag	attgtgtgcc	caaggtccag	tgtccctggt	actatgatgg	2340
tgaagctctt	cagcccggaag	acatcttctc	agaccatcac	accatgtgct	actgtgagga	2400
tggcttcctg	cactgtacca	caagtggagg	cctgggaagg	ctgtctccca	acccgggtgt	2460
cagcagccccc	cggtgtcacc	gcagcaaaag	gagcctgtcc	tgtccggccc	ccatgggtcaa	2520
gttgggtggt	cccgctgata	acccgagggc	tgaaggactg	gagtggtgca	aaacttgcca	2580
gaactatgat	ctgcagtgca	tgagcacagg	ctgtgtctcc	ggtgtccctc	gcccggaggg	2640
catggtccgg	catgaaaaaca	ggtgtgtggc	gtggaaaaga	tgtccctgtc	tccacccagg	2700
ccaaagagtar	gcctcaggag	aaactgtgaa	aattgaactg	aaacttctgt	tctgtcggga	2760
ccggaagtgg	acctgcacag	accatgtgtg	tgatcccact	tgtctcgcca	tgggcctggc	2820
gcactacctc	accttcgacg	gaactcaagta	cctgttccct	ggggagtgcc	agtatgtctc	2880
ggtgcaggat	tactcgggca	gtaacccctg	gacettaagg	atcctgggtg	ggaaacgagg	2940
gtgcagctac	ccctcagtg	aatgcaagaa	gcccgtcacc	atcctgggtg	aaggaggaga	3000
gattgaactg	tttgatgggg	aggtgaatgt	gaagaaaacc	atgaaggatg	agactcactt	3060
tgaggtggta	gagtgctggt	agtacgtcat	tctgtctctg	ggcaaggcac	tctctgtggt	3120
ctggggaacc	cgcctgagca	tctctgtgac	cctgaaggcg	acataccagg	aggaggtgtg	3180
tggcctgtgt	gggaattttg	atggcctcca	gaacaatgat	ttcaccagca	gcagcctcca	3240
aatagaagaa	gacctgtggg	actttgggaa	ttcctggaaa	gtgaaacccg	agtgtgcgca	3300
caccaagaaa	gtaccactgg	actcatcccc	tgcctgtctg	cacaaacaaca	tcatgaagca	3360
gacgatgggt	gattctctct	gcaggatcct	caccagtgat	atlttccagg	actgcaacag	3420
gctggtggag	cctgagccat	tcttgacat	ttgcattctac	gacacttgcct	cctgtgagtc	3480
cattggggac	tgcactgtct	tctgtgacac	cattgtctgt	tacggccacg	tctgtgccc	3540
gcctggcaag	gtggtagcct	ggaggacccg	cacattctct	ccccagaatt	gcgaggagcg	3600
gaattctcac	gaggaatgggt	atgagtggtg	gtggccttat	aaagctgtgt	ccctgcctg	3660
tcccatcagg	tgcacgcacc	ccgagccact	ggatggaat	gtacagtggt	ttgaagggtg	3720
ccatgctaac	tccctccag	gaaaaatcct	ggtcaggtt	ttcagagact	gcctcgaccc	3780
tgaagactgt	cctgtgtgag	agggtggctg	tctcgggtt	ttcagagaa	agaaaatcat	3840
cttgaacccc	agtgaacctg	agcactgcca	aatgtgtaat	tgtgtgtgtg	tcaacttcac	3900
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	3960
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4020
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4080
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4140
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4200
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4260
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4320
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4380
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4440
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4500
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4560
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4620
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4680
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4740
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4800
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4860
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4920
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	4980
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5040
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5100
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5160
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5220
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5280
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5340
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5400
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5460
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5520
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5580
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5640
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5700
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5760
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5820
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5880
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	5940
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6000
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6060
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6120
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6180
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6240
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6300
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6360
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6420
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6480
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6540
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6600
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6660
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6720
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6780
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6840
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6900
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	6960
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7020
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7080
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7140
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7200
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7260
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7320
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7380
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7440
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7500
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7560
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7620
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7680
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7740
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7800
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7860
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7920
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	7980
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8040
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8100
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8160
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8220
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8280
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8340
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8400
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8460
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8520
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8580
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8640
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8700
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8760
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8820
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8880
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	8940
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9000
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9060
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9120
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9180
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9240
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9300
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9360
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9420
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9480
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9540
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9600
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9660
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9720
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9780
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9840
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9900
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	9960
ctctggtggt	tcaatgagac	cagggaagtt	tctgtgtgt	ctcagatgtg	gtccctattc	10020

ctctggtggt tcaatgagac cagggaagtt tctgtgtgt ctcagatgtg gtccctattc 10080

ctgtctggccc acccagacgg agcccatgca ggtgggcccgtg cgttgccacca atggtctccct 8580
 catctaccat gagatccctca atgccatgca atgcagggtgt tccccccagga agtgccagcaa 8640
 gtgaaggccac tgcctggatg ctactgtgca ctgccttaac cgacctcaact ggaatgggcca 8700
 gaggctctgt cagtcctcct cagtcctcct cctgctctga tcttggtgctt cctgatccca 8760
 caataaaggt caatctttca ccttgaaaaa aaaaaaaaaa aa 8802

<210> 2

<211> 2813

<212> PRT

<213> Canis familiaris

<100> 2

Met Ser Pro Thr Arg Leu Val Arg Val Leu Leu Ala Leu Ala Leu Ile
 1 5 10 15

Leu Pro Gly Lys Leu Cys Thr Lys Gly Thr Val Gly Arg Ser Ser Met
 20 25 30

Ala Arg Cys Ser Leu Leu Gly Gly Asp Phe Ile Asn Thr Phe Asp Glu
 35 40 45

Ser Met Tyr Ser Phe Ala Gly Asp Cys Ser Tyr Leu Leu Ala Gly Asp
 50 55 60

Cys Gln Glu His Ser Ile Ser Leu Ile Gly Gly Phe Gln Asn Asp Lys
 65 70 75 80

Arg Val Ser Leu Ser Val Tyr Leu Gly Glu Phe Phe Asp Ile His Leu
 85 90 95

Phe Val Asn Gly Thr Met Leu Gln Gly Thr Gln Ser Ile Ser Met Pro
 100 105 110

Tyr Ala Ser Asn Gly Leu Tyr Leu Glu Ala Glu Ala Gly Tyr Tyr Lys
 115 120 125

Leu Ser Ser Glu Ala Tyr Gly Phe Val Ala Arg Ile Asp Gly Asn Gly
 130 135 140

Asn Ile Gln Tyr Leu Leu Ser Asp Ala Tyr Ile Asn Lys Thr Tyr Gly
 145 150 155 160

Ser Cys Gly Asn Phe Asn Ile Ile Ala Gln Asp Asp Phe Lys Thr Gln
 165 170 175

Glu Gly Thr Leu Thr Ser Asp Pro Tyr Asp Phe Ala Asn Ser Trp Ala
 180 185 190

Ser Ser Gly Gly Gln Gln Asn Tyr Lys Asp Ala Val Ser Ile Ile Ser Ser
 195 200 205 210 215 220

Val Asp Pro Glu Pro Phe Val Ala Leu Cys Glu Arg Thr Leu Cys Thr
245 250 255

Cys Val Gln Gly Met Glu Cys Pro Cys Ala Val Leu Leu Glu Tyr Ala
260 265 270

Arg Ala Cys Ala Gln Gln Gly Ile Val Leu Tyr Gly Trp Thr Asp His
275 280 285

Ser Val Cys Arg Pro Ala Cys Pro Ala Gly Met Glu Tyr Lys Glu Cys
290 295 300

Val Ser Pro Cys Thr Arg Thr Cys Gln Ser Leu His Val Lys Glu Val
305 310 315 320

Cys Gln Glu Gln Cys Val Asp Gly Cys Ser Cys Pro Glu Gly Gln Leu
325 330 335

Leu Asp Glu Gly His Cys Val Gly Ser Ala Glu Cys Ser Cys Val His
340 345 350

Ala Gly Gln Arg Tyr Pro Pro Gly Ala Ser Leu Leu Gln Asp Cys His
355 360 365

Thr Cys Ile Cys Arg Asn Ser Leu Trp Ile Cys Ser Asn Glu Glu Cys
370 375 380

Pro Gly Glu Cys Leu Val Thr Gly Gln Ser His Phe Lys Ser Phe Asp
385 390 395 400

Asn Arg Tyr Phe Thr Phe Ser Gly Val Cys His Tyr Leu Leu Ala Gln
405 410 415

Asp Cys Gln Asp His Thr Phe Ser Val Val Ile Glu Thr Val Gln Cys
420 425 430

Ala Asp Asp Leu Asp Ala Val Cys Thr Arg Ser Val Thr Val Arg Leu
435 440 445

Pro Gly His His Asn Ser Ile Val Lys Leu Lys Asn Lys Lys Lys Val
445 450 455

Ser Met Asp Gly His Asp Ile Gln Ile Trp Leu Leu His His Arg Leu
455 460 465 470

Arg Ile Gln His Thr Val Met Ala Ser Val Arg Leu Ser Tyr Gly His
475 480 485

Asn Leu Thr Met Asp Ser Asp Thr Asn Lys Asn Leu Leu Val Thr Leu
490 495

Leu Val Glu Asp Phe Gly Asn Ala Trp Lys Leu Leu Gly Ala Cys Glu
545 550 555 560

Asn Leu Gln Lys Gln His Arg Asp Pro Cys Ser Leu Asn Pro Arg Gln
565 570 575

Ala Arg Phe Ala Glu Glu Ala Cys Ala Leu Leu Thr Ser Ser Lys Phe
580 585 590

Glu Pro Cys His Arg Ala Val Gly Pro Gln Pro Tyr Val Gln Asn Cys
595 600 605

Leu Tyr Asp Val Cys Ser Cys Ser Asp Gly Arg Asp Cys Leu Cys Ser
610 615 620

Ala Val Ala Asn Tyr Ala Ala Ala Val Ala Arg Arg Gly Val His Ile
625 630 635 640

Ala Trp Arg Glu Pro Gly Phe Cys Ala Leu Ser Cys Pro Gln Gly Gln
645 650 655

Val Tyr Leu Gln Cys Gly Thr Pro Cys Asn Met Thr Cys Leu Ser Leu
660 665 670

Ser Tyr Pro Glu Glu Asp Cys Asn Glu Val Cys Leu Glu Ser Cys Phe
675 680 685

Ser Pro Pro Gly Leu Tyr Leu Asp Glu Arg Gly Asp Cys Val Pro Lys
690 695 700

Ala Gln Cys Pro Cys Tyr Tyr Asp Gly Glu Ile Phe Gln Pro Glu Asp
705 710 715 720

Ile Phe Ser Asp His His Thr Met Cys Tyr Cys Glu Asp Gly Phe Met
725 730 735

His Cys Thr Thr Ser Gly Gly Leu Gly Ser Leu Leu Pro Asn Pro Val
740 745 750

Leu Leu Ser Leu Asn Tyr His Asn Ser Lys Asn Ser Leu Ser Tyr Asn
755 760 765 770

Ala Leu Met Val Lys Leu Val Tyr Leu Ala Asp Asn Leu Asn Ala Gln
775 780 785 790

Gly Leu Ala Cys Ala Lys Thr Cys Gln Asn Tyr Asp Leu Gln Cys Met
795 800 805 810

Pro Leu Gly Tyr Val Ser Gly Cys Leu Tyr Leu Leu Tyr Met Val Asn
815 820 825 830

Cys Val Cys Arg Asp Arg Lys Trp Thr Cys Thr Asp His Val Cys Asp
850 855 860

Ala Thr Cys Ser Ala Ile Gly Met Ala His Tyr Leu Thr Phe Asp Gly
865 870 875 880

Leu Lys Tyr Leu Phe Pro Gly Glu Cys Gln Tyr Val Leu Val Gln Asp
885 890 895

Tyr Cys Gly Ser Asn Pro Gly Thr Leu Arg Ile Leu Val Gly Asn Glu
900 905 910

Gly Cys Ser Tyr Pro Ser Val Lys Cys Lys Lys Arg Val Thr Ile Leu
915 920 925

Val Glu Gly Gly Glu Ile Glu Leu Phe Asp Gly Glu Val Asn Val Lys
930 935 940

Lys Pro Met Lys Asp Glu Thr His Phe Glu Val Val Glu Ser Gly Gln
945 950 955 960

Tyr Val Ile Leu Leu Leu Gly Lys Ala Leu Ser Val Val Trp Asp His
965 970 975

Arg Leu Ser Ile Ser Val Thr Leu Lys Arg Thr Tyr Gln Glu Gln Val
980 985 990

Cys Gly Leu Cys Gly Asn Phe Asp Gly Ile Gln Asn Asn Asp Phe Thr
995 1000 1005

Ser Ser Ser Leu Gln Ile Glu Glu Asp Pro Val Asp Phe Gly Asn Ser
1010 1015 1020

Trp Lys Val Asn Pro Gln Cys Ala Asp Thr Lys Lys Val Pro Leu Asp
1025 1030 1035 1040

Ser Ser Pro Ala Val Cys His Asn Asn Ile Met Lys Gln Thr Met Val
1045 1050 1055

Arg Ser Ser Cys Arg Ile Leu Thr Ser Arg Ile Ser Ile Arg Tyr Asn
1060 1065 1070

Arg Leu Val Arg Ile Gln Ile Thr Leu Asp Ile Tyr Ile Tyr Arg Thr
1075 1080 1085

Cys Ser Cys Gln Ser Ile Gly Asp Cys Thr Cys Ile Cys Asp Thr Ile
1090 1095 1100

Ala Ala Tyr Arg His Tyr Tyr Ala Gln His Gly Lys Val Val Ala Ile
1105 1110

Cys Pro Ile Thr Cys Gln His Pro Glu Pro Leu Ala Cys Pro Val Gln
1155 1160 1165

Cys Val Glu Gly Cys His Ala His Cys Pro Pro Gly Lys Ile Leu Asp
1170 1175 1180

Glu Leu Leu Gln Thr Cys Ile Asp Pro Glu Asp Cys Pro Val Cys Glu
1185 1190 1195 1200

Val Ala Gly Arg Arg Leu Ala Pro Gly Lys Lys Ile Ile Leu Asn Pro
1205 1210 1215

Ser Asp Pro Glu His Cys Gln Ile Cys Asn Cys Asp Gly Val Asn Phe
1220 1225 1230

Thr Cys Lys Ala Cys Arg Glu Pro Gly Ser Val Val Val Pro Pro Thr
1235 1240 1245

Asp Gly Pro Ile Gly Ser Thr Thr Ser Tyr Val Glu Asp Thr Ser Glu
1250 1255 1260

Pro Pro Leu His Asp Phe His Cys Ser Arg Leu Leu Asp Leu Val Phe
1265 1270 1275 1280

Leu Leu Asp Gly Ser Ser Lys Leu Ser Glu Asp Glu Phe Glu Val Leu
1285 1290 1295

Lys Val Phe Val Val Gly Met Met Glu His Leu His Ile Ser Gln Lys
1300 1305 1310

Arg Ile Arg Val Ala Val Val Glu Tyr His Asp Gly Ser His Ala Tyr
1315 1320 1325

Ile Glu Leu Lys Asp Arg Lys Arg Pro Ser Glu Leu Arg Arg Ile Thr
1330 1335 1340

Ser Gln Val Lys Tyr Ala Gly Ser Glu Val Ala Ser Thr Ser Glu Val
1345 1350 1355 1360

Leu Lys Tyr Thr Leu Ile Ala Ile His Gly Lys Ile Arg Arg Ile Thr
1365 1370 1375 1380

Ala Ser Arg Ile Ala Ile Ile Ile Met Ala Ser His His Thr Ser Arg
1385 1390 1395 1400

Leu Ala Arg Asn Leu Val Arg Tyr Val His Gly Leu Lys Lys Lys Lys
1405 1410 1415 1420

Val Ile Val Ile Ile Val Lys Ile His Ile His Ala Ser Leu Lys Ile
1425 1430 1435 1440

Leu Cys Asp Leu Ala Pro Glu Ala Pro Ala Pro Thr Gln His Pro Pro
1460 1465 1470

Met Ala Gln Val Thr Val Gly Ser Glu Leu Leu Gly Val Ser Ser Pro
1475 1480 1485

Gly Pro Lys Arg Asn Ser Met Val Leu Asp Val Val Phe Val Leu Glu
1490 1495 1500

Gly Ser Asp Lys Ile Gly Glu Ala Asn Phe Asn Lys Ser Arg Glu Phe
1505 1510 1515 1520

Met Glu Glu Val Ile Gln Arg Met Asp Val Gly Gln Asp Arg Ile His
1525 1530 1535

Val Thr Val Leu Gln Tyr Ser Tyr Met Val Thr Val Glu Tyr Thr Phe
1540 1545 1550

Ser Glu Ala Gln Ser Lys Gly Glu Val Leu Gln Gln Val Arg Asp Ile
1555 1560 1565

Arg Tyr Arg Gly Gly Asn Arg Thr Asn Thr Gly Leu Ala Leu Gln Tyr
1570 1575 1580

Leu Ser Glu His Ser Phe Ser Val Ser Gln Gly Asp Arg Glu Gln Val
1585 1590 1595 1600

Pro Asn Leu Val Tyr Met Val Thr Gly Asn Pro Ala Ser Asp Glu Ile
1605 1610 1615

Lys Arg Met Pro Gly Asp Ile Gln Val Val Pro Ile Gly Val Gly Pro
1620 1625 1630

His Ala Asn Val Gln Glu Leu Glu Lys Ile Gly Trp Pro Asn Ala Pro
1635 1640 1645

Ile Leu Ile His Asp Phe Glu Met Leu Pro Arg Glu Ala Pro Asp Leu
1650 1655 1660

Glu Leu Ala Asn Tyr Tyr Ser Gly Glu Gly Ser Gln Ile Leu Thr Leu
1665 1670 1675 1680

Arg Pro Thr Leu Asp Tyr Ser Gln Pro Leu Asp Thr Val Leu Ile Leu
1685 1690 1695

Asp Gly Ser Ser Ser Ile Pro Ala Ser Tyr Phe Asp Glu Met Lys Ser
1700 1705 1710

Phe Thr Lys Ala Ile Leu Ser Asn Ala Asn Ile Gly Leu Asn Leu Ile
1715 1720 1725

Met Gln Gln Gln Gly Gly Pro Ser Glu Ile Gly Asp Ala Leu Ser Phe
1765 1770 1775

Ala Val Arg Tyr Val Thr Ser Glu Val His Gly Ala Arg Pro Gly Ala
1780 1785 1790

Ser Lys Ala Val Val Ile Leu Val Thr Asp Val Ser Val Asp Ser Val
1795 1800 1805

Asp Ala Ala Ala Glu Ala Ala Arg Ser Asn Arg Val Thr Val Phe Pro
1810 1815 1820

Ile Gly Ile Gly Asp Arg Tyr Ser Glu Ala Gln Leu Ser Ser Leu Ala
1825 1830 1835 1840

Gly Pro Lys Ala Gly Ser Asn Met Val Arg Leu Gln Arg Ile Glu Asp
1845 1850 1855

Leu Pro Thr Val Ala Thr Leu Gly Asn Ser Phe Phe His Lys Leu Cys
1860 1865 1870

Ser Gly Phe Asp Arg Val Cys Val Asp Glu Asp Gly Asn Glu Lys Arg
1875 1880 1885

Pro Gly Asp Val Trp Thr Leu Pro Asp Gln Cys His Thr Val Thr Cys
1390 1895 1900

Leu Pro Asp Gly Gln Thr Leu Leu Lys Ser His Arg Val Asn Cys Asp
1905 1910 1915 1920

Arg Gly Pro Arg Pro Ser Cys Pro Asn Gly Gln Pro Pro Leu Arg Val
1925 1930 1935

Glu Glu Thr Cys Gly Cys Arg Trp Thr Cys Pro Cys Val Cys Met Gly
1940 1945 1950

Ser Ser Thr Arg His Ile Val Thr Phe Asp Gly Gln Asn Phe Lys Leu
1955 1960 1965

His Gly Ser Tyr Ser Tyr Val Ser His His Asn Lys Ser Ser Arg Leu
1965 1970 1975

His Val Ile Ser His Asn Gly Ala Tyr Ser Phe Gly Ala Lys His Thr
1975 1980 1985 1990

Cys Met Lys Ser Ile Glu Val Lys His Asp Gly Leu Ser Val Glu Leu
1985 1990 1995

His Ser Arg Ser Ser Ser Thr Val Asn Gly Arg Leu Val Ser Leu Leu

Asn Asn Glu Phe Gln Leu Gln Leu Ser Pro Arg Thr Phe Ala Ser Lys
2065 2070 2075 2080

Thr Tyr Gly Leu Cys Gly Ile Cys Asp Glu Asn Gly Ala Asn Asp Phe
2085 2090 2095

Ile Leu Arg Asp Gly Thr Val Thr Thr Asp Trp Lys Ala Leu Ile Gln
2100 2105 2110

Glu Trp Thr Val Gln Gln Leu Gly Lys Thr Ser Gln Pro Val His Glu
2115 2120 2125

Glu Gln Cys Pro Val Ser Glu Phe Phe His Cys Gln Val Leu Leu Ser
2130 2135 2140

Glu Leu Phe Ala Glu Cys His Lys Val Leu Ala Pro Ala Thr Phe Tyr
2145 2150 2155 2160

Ala Met Cys Gln Pro Asp Ser Cys His Pro Lys Lys Val Cys Glu Ala
2165 2170 2175

Ile Ala Leu Tyr Ala His Leu Cys Arg Thr Lys Gly Val Cys Val Asp
2180 2185 2190

Trp Arg Arg Ala Asn Phe Cys Ala Met Ser Cys Pro Pro Ser Leu Val
2195 2200 2205

Tyr Asn His Cys Glu His Gly Cys Pro Arg Leu Cys Glu Gly Asn Thr
2210 2215 2220

Ser Ser Cys Gly Asp Gln Pro Ser Glu Gly Cys Phe Cys Pro Pro Asn
2225 2230 2235 2240

Gln Val Met Leu Glu Gly Ser Cys Val Pro Glu Glu Ala Cys Thr Gln
2245 2250 2255

Cys Ile Ser Glu Asp Gly Val Arg His Gln Phe Leu Glu Thr Trp Val
2260 2265 2270

Ile Ala His Ser Ile Tyr Leu Ile Tyr Thr Tyr Ile Ser Gly Asn Lys
2275 2280 2285 2290

Val Asn Tyr Thr Ile Leu Ile Tyr Ile Thr Ala Tyr Ala Ser Thr Tyr
2295 2300 2305 2310

Gly Pro Cys Glu Val Ala Arg Leu Arg Gln Asn Ala Val Gln Tyr Cys
2315 2320 2325 2330

Ile Gln Tyr Ser Tyr Val Ser Arg Ser Val Ser Tyr Asn Ile Ile Ile

Arg Glu Ser Pro Pro Ser Cys Pro Pro His Arg Thr Pro Ala Leu Arg
2370 2375 2379

Lys Thr Gln Cys Cys Asp Glu Tyr Glu Cys Ala Cys Asn Cys Val Asn
2385 2390 2395 2400

Ser Thr Val Ser Cys Pro Leu Gly Tyr Leu Ala Ser Ala Val Thr Asn
2405 2410 2415

Asp Cys Gly Cys Thr Thr Thr Thr Cys Phe Pro Asp Lys Val Cys Val
2420 2425 2430

His Arg Gly Thr Ile Tyr Pro Val Gly Gln Phe Trp Glu Glu Ala Cys
2435 2440 2445

Asp Val Cys Thr Cys Thr Asp Leu Glu Asp Ser Val Met Gly Leu Arg
2450 2455 2460

Val Ala Gln Cys Ser Gln Lys Pro Cys Glu Asp Asn Cys Leu Ser Gly
2465 2470 2475 2480

Phe Thr Tyr Val Leu His Glu Gly Glu Cys Cys Gly Arg Cys Leu Pro
2485 2490 2495

Ser Ala Cys Glu Val Val Thr Gly Ser Pro Arg Gly Asp Ala Gln Ser
2500 2505 2510

His Trp Lys Asn Val Gly Ser His Trp Ala Ser Pro Asp Asn Pro Cys
2515 2520 2525

Leu Ile Asn Glu Cys Val Arg Val Lys Glu Glu Val Phe Val Gln Gln
2530 2535 2540

Arg Asn Val Ser Cys Pro Gln Leu Asn Val Pro Thr Cys Pro Thr Gly
2545 2550 2555 2560

Phe Gln Leu Ser Cys Lys Thr Ser Glu Cys Cys Pro Thr Cys His Cys
2565 2570 2575

His Ile Leu Ala Tyr Leu Leu Asn Gly Thr Leu Ile Gly Ile Gly
2580 2585 2590 2595

Lys Pro Leu Met Ile Asp Val Tyr Thr Thr Tyr Asn Tyr Thr Val Ile
2600 2605 2610

Val Gly Val Ile Ser Gly Phe Lys Leu Glu Gly Arg Lys Thr Thr Cys
2615 2620 2625

His Ala Tyr Ile Leu Gly Tyr Tyr Thr His Lys Asn Ile Gly Ile Tyr

Ser His Phe Cys Lys Val Asn Glu Arg Gly Glu Tyr Ile Trp Glu Lys
2675 2680 2685

Arg Val Thr Gly Cys Pro Pro Phe Asp Glu His Lys Cys Leu Ala Glu
2690 2695 2700

Gly Gly Lys Ile Met Lys Ile Pro Gly Thr Cys Cys Asp Thr Cys Glu
2705 2710 2715 2720

Glu Pro Glu Cys Lys Asp Ile Ile Ala Lys Leu Gln Arg Val Lys Val
2725 2730 2735

Gly Asp Cys Lys Ser Glu Glu Glu Val Asp Ile His Tyr Cys Glu Gly
2740 2745 2750

Lys Cys Ala Ser Lys Ala Val Tyr Ser Ile His Met Glu Asp Val Gln
2755 2760 2765

Asp Gln Cys Ser Cys Cys Ser Pro Thr Gln Thr Glu Pro Met Gln Val
2770 2775 2780

Ala Leu Arg Cys Thr Asn Gly Ser Leu Ile Tyr His Glu Ile Leu Asn
2785 2790 2795 2800

Ala Ile Glu Cys Arg Cys Ser Pro Arg Lys Cys Ser Lys
2805 2810

<210> 3

<211> 60

<212> DNA

<213> Canis familiaris

<400> 3

ATTGTTTCTG CAGGATGCTG GAGGATGAG CAGGATGAG TATGATGAG ATTGTTTCTG

<210> 4

<211> 60

<212> DNA

<213> Canis familiaris

<400> 4

ATTGTTTCTG CAGGATGCTG GAGGATGAG CAGGATGAG TATGATGAG ATTGTTTCTG

<210> 5

<211> 60

<212> DNA

<213> Canis familiaris

<400> 6
gagggagagc gggccacagc acacgagga aatgtgcccaggaaagtga gggcaggac 60

<210> 7
<211> 60
<212> DNA
<213> Canis familiaris

<400> 7
tgggtgaaag ccccatatcc cgaactcttg tcaaggagac ttctaccacaggccacagac 60

<210> 8
<211> 60
<212> DNA
<213> Canis familiaris

<400> 8
ctggagcagc ggtctggggc tggcaggctg agggacatgg aggaatgca tgagaagcac 60

<210> 9
<211> 58
<212> DNA
<213> Canis familiaris

<400> 9
gttctctgag ctctctcttg tcccaccagc atctccatgc cctacgcctc caatgggc 58

<210> 10
<211> 24
<212> DNA
<213> Canis familiaris

<400> 10
aatgacaaa agagtggagc ggta 24

<210> 11
<211> 24
<212> DNA
<213> Canis familiaris

<400> 11
ggtgctgctg cctgctgctg cctgctgctg cctgctgctg cctgctgctg 24

<210> 12
<211> 30
<212> PBT
<213> Canis familiaris

<400> 12
ggtgctgctg cctgctgctg cctgctgctg cctgctgctg cctgctgctg

<210> 13
<211> 9
<212> PRT
<213> Canis familiaris

<400> 13
Ile Ser Met Pro Tyr Ala Ser Asn Gly
1 5

<210> 14
<211> 31
<212> DNA
<213> Canis familiaris

<400> 14
aggacaactg cctgcctgtc ggtgagtggg g 31

<210> 15
<211> 31
<212> DNA
<213> Canis familiaris

<400> 15
aggacaactg cctgcctgtc agtgagtggg g 31

<210> 16
<211> 8
<212> DNA
<213> Canis familiaris

<400> 16
aggtaag 8

<210> 17
<211> 12
<212> DNA
<213> Canis familiaris

<400> 17
tttttttttt 12

<210> 18
<211> 17
<212> DNA
<213> Canis familiaris

<400> 19
tgggttggc agccc 19

<210> 20
<211> 16
<212> DNA
<213> Canis familiaris

<400> 20
gaggttttg tggccc 16

<210> 21
<211> 218
<212> DNA
<213> Canis familiaris

<400> 21
gtcctgtggg agcagtgccc gctcctgaag agtgcctcgg tgtttgcccg ctgccacccg 60
ctggtggacc ctgagccttt tctcgccttg tctgaaagga ctctgtgcac ctgtgtccag 120
gggatggagt gcccttgtgc ggtcctcctg gactacgccc gggcctgtgc ccagcagggg 180
attgtgtgt acggtggac cgaccacagc gtctggc 218

<210> 22
<211> 73
<212> PRT
<213> Canis familiaris

<400> 22
Val Leu Trp Glu Gln Cys Gln Leu Leu Lys Ser Ala Ser Val Phe Ala
1 5 10 15

Arg Cys His Pro Leu Val Asp Pro Gln Pro Ile Val Ala Leu Cys Glu
20 25 30

Arg Thr Leu Cys Thr Cys Val Gln Gly Met Gln Cys Pro Cys Ala Val
35 40 45

Leu Leu Glu Tyr Ala Arg Ala Cys Ala Gln Gln Gly Ile Val Leu Tyr
50 55 60 65 70 75

Gly Thr Thr Arg His Arg Thr Tyr Arg

<210> 23
<211> 73
<212> DNA
<213> Canis familiaris

<223> n a, c, r, or d

<400> 24

qennnnnnn

11

 $\langle 210 \rangle \rightarrow 2^5$

211 > 18

(212) DNA

<213> Canis familiaris

<400> 25

gtggtaggtc caqccgta

18

<210> 26

<211> 20

<212> DNA

(213) Canis familiaris

<400>: 26

tctaccctgt gggccagttc

20

<210> 27

<221> 16

<214> DIA

<213> Canis familiaris

435 27

gacvanon: 2011-01-01

2203 23

4223 21

62 223

623 - Girls' Athletics

110 2

$\chi^2 = 1.0$ $df = 1$ $p = 0.32$ $\phi^2 = 0.00$ $95\% CI = [-0.01, 0.01]$ $OR = 1.0$ $95\% CI = [0.4, 2.4]$

1. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Arar and Collins (1971) using a Shimadzu 1010 spectrophotometer.

[illegible]

1. *Chlorophyll a* (Chl *a*)

[illegible]